

AMENDMENTS TO THE CLAIMS

For the Examiner's convenience, Applicants provide the following listing of the claims:

1. – 19. (Canceled)

20. (Currently Amended) A glutamic acid synthesizing gene selected from the group consisting of glutamate dehydrogenase, citrate synthase, isocitrate ~~synthase~~ dehydrogenase, pyruvate dehydrogenase, and aconitase, comprising a DNA sequence situated at position -35 in a promoter sequence of the glutamic acid synthesizing gene, wherein said DNA sequence is selected from the group consisting of CGGTCA, TTGTCA, TTGACA, and TTGCCA.

21. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, which is glutamate dehydrogenase.

22. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, which is citrate synthase.

23. (Currently Amended) The glutamic acid synthesizing gene of Claim 20, which is isocitrate ~~synthase~~ dehydrogenase.

24. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, which is pyruvate dehydrogenase.

25. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, which is aconitase.

26. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, further comprising TATAAT or CATAAT situated at position -10 in the promoter sequence.

27. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, wherein said DNA sequence is CGGTCA.

28. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, wherein said DNA sequence is TTGTCA.

29. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, wherein said DNA sequence is TTGACA.

30. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, wherein said DNA sequence is TTGCCA.

31. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, which is citrate synthase and wherein said DNA sequence is TTGACA.

32. (Previously Presented) The glutamic acid synthesizing gene of Claim 31, further comprising TATAAT situated at position -10 in the promoter sequence.

33. (Currently Amended) The glutamic acid synthesizing gene of Claim 20, which is isocitrate ~~synthase~~ dehydrogenase and wherein said DNA sequence is TTGCCA or TTGACA.

34. (Previously Presented) The glutamic acid synthesizing gene of Claim 33, further comprising TATAAT situated at position -10 in the promoter sequence.

35. (Previously Presented) The glutamic acid synthesizing gene of Claim 20, which is pyruvate dehydrogenase and wherein said DNA sequence is TTGCCA.

36. (Previously Presented) The glutamic acid synthesizing gene of Claim 35, further comprising TATAAT.

37. (Previously Presented) A coryneform bacterium comprising the glutamic acid synthesizing gene of Claim 20.